CDPK2 (Genbank accession D21806)

Amino Acid Sequence (SEQ ID NO: 1)

METKPNPRRP	SNTVLPYQTP	RLRDHYLLGK	KLGQGQFGTT	YLCTEKSTSA	NYACKSIPKR
KLVCREDYED	VWREIQIMHH	LSEHPNVVRI	KGTYEDSVFV	HIVMEVCEGG	ELFDRIVSKG
HFSEREAVKL	IKTILGVVEA	CHSLGVMHRD	LKPENFLFDS	PKDDAKLKAT	DFGLSVFYKP
GOYLYDVVGS	PYYVAPEVLK	KCYGPEIDVW	SAGVILYILL	SGVPPFWAET	ESGIFRQILQ
GKLDFKSDPW	PTISEAAKDL	IYKMLERSPK	KRISAHEALC	HPWIVDEQAA	PDKPLDPAVL
SRLKQFSQMN	KIKKMALRVI	AERLSEEEIG	GLKELFKMID	TDNSGTITFE	ELKAGLKRVG
SELMESEIKS	LMDAADIDNS	GTIDYGEFLA	ATLHMNKMER	EEILVAAFSD	FDKDGSGYIT
IDELOSACTE	FGLCDTPLDD	MIKEIDLDND	GKIDFSEFTA	MMRKGDGVGR	SRTMMKNLNF
NIADAFGVDG	EKSDD				

Nucleotide Sequence (SEQ ID NO: 2)

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     acaccacgat taagagatca ttaccttctg ggaaaaaagc taggccaagg ccaatttgga 181
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CDPK4 (Genbank accession number U31752)

Amino Acid Sequence (SEQ ID NO: 3)

MEKPNPRRPS	NSVLPYETPR	LRDHYLLGKK	LGQGQFGTTY	LCTEKSSSAN	YACKSIPKRK
LVCREDYEDV	WREIQIMHHL	SEHPNVVRIK	GTYEDSVFVH	IVMEVCEGGE	LFDRIVSKGC
FSEREAAKLI	KTILGVVEAC	HSLGVMHRDL	KPENFLFDSP	SDDAKLKATD	FGLSVFYKPG
QYLYDVVGSP	YYVAPEVLKK	CYGPEIDVWS	AGVILYILLS	GVPPFWAETE	SGIFRQILQG
KIDFKSDPWP	TISEGAKDLI	YKMLDRSPKK	RISAHEALCH	PWIVDEHAAP	DKPLDPAVLS
RLKQFSQMNK	IKKMALRVIA	ERLSEEEIGG	LKELFKMIDT	DNSGTITFEE	LKAGLKRVGS
ELMESEIKSL	MDAADIDNSG	TIDYGEFLAA	TLHINKMERE	ENLVVAFSYF	DKDGSGYITI
DELQQACTEF	GLCDTPLDDM	IKEIDLDNDG	KIDFSEFTAM	MKKGDGVGRS	RTMRNNLNFN
IAEAFGVEDT	SSTAKSDDSP	K			

Nucleotide Sequence (SEQ ID NO: 4)

1 atqqagaaac caaaccctag aagaccctca aacagtgttc ttccatacga aacaccaaga 61 ttaagagatc actatctcct cggcaaaaag ctaggccaag gccaatttgg aacaacctat 121 ctctqtacag agaaatcatc atcagctaat tacqcttgca aatcaatccc aaaacgtaag 181 cttgtatqtc gtgaagacta cgaagatgta tggcgtgaga ttcagatcat gcatcatctc 241 totgagoato otaatgttqt tagaatcaag ggtacttatg aagactotgt ttttgttcac 301 attgttatgg aagtttgtga aggtggtgag ctttttgatc ggattgtttc taaagggtgt 361 tttagtgaac gtgaagctgc taagttgatt aagactattc ttggtgttgt tgaggcttgt 421 cattetettg gtgttatgca tagagatett aageetgaga atttettgtt tgatagteec 481 agtgatgatg ctaagcttaa agctacagac tttggtttgt ctgtcttcta caagccaggg 541 caqtatctqt atqatqtaqt tqqaaqtccq tattatqttq cacctgaggt tctgaagaaa 601 tgttatggac cagagataga cgtgtggagc gccggtgtta tcttgtacat cttactaagt 661 ggggttcctc ctttttgggc agaaaccgag tcaggaatct ttaggcagat attgcaaggg 721 aagatagatt ttaaatctga tccgtggcct actatctcag aaggtgctaa agatttgatt 781 tacaaaatqc tcqatagqag ccccaagaaa cgtatttctg cacatgaagc attgtgtcac 841 ccttggattg ttgatgaaca tgctgcacca gacaagcctc tcgacccagc agtcttgtcg 901 cqacttaagc agttctcqca aatgaataaa atcaaqaaaa tqqccttacq aqtaatcqcg 961 gagagactct cggaggaaga gattggtggt ctgaaggaat tgttcaaaat gatagataca 1021 gacaacagtg gaacaatcac ctttgaagag cttaaagcag gtctaaagag agttggatct 1081 gaattgatgg aatcagaaat caagtctctt atggatgcgg cggatataga caacagtggg 1141 acaatagact acggtgaatt cctagcagcg acattacata taaacaagat ggagagagaa 1201 gagaacttgg tggttgcgtt ttcatacttt gataaagatg gtagcggtta tatcaccatt 1261 gacgagette aacaageetg cacagagttt ggtetetgtg acaeteetet tgatgacatg 1321 atcaaaqaqa ttqatcttqa taatgacqqq aagattgatt tctcagagtt tactgctatg 1381 atgaagaaag gagatggtgt tgggaggagc agaactatga ggaacaactt gaacttcaat 1441 atagctgaág ettttggagt tgaggacaca agcagcactg etaaatetga tgattcacca 1501 aagtaattat aatcatctat atacttggaa ttgagaaatg agaactcaca aaagaaaaac 1561 tgaatctttc ctttttgttt tcgtttccac ttttgtagat gagcaacttt ctcaattttg 1621 ttataagcat ggataatttt gcttcatatt ttctgcg